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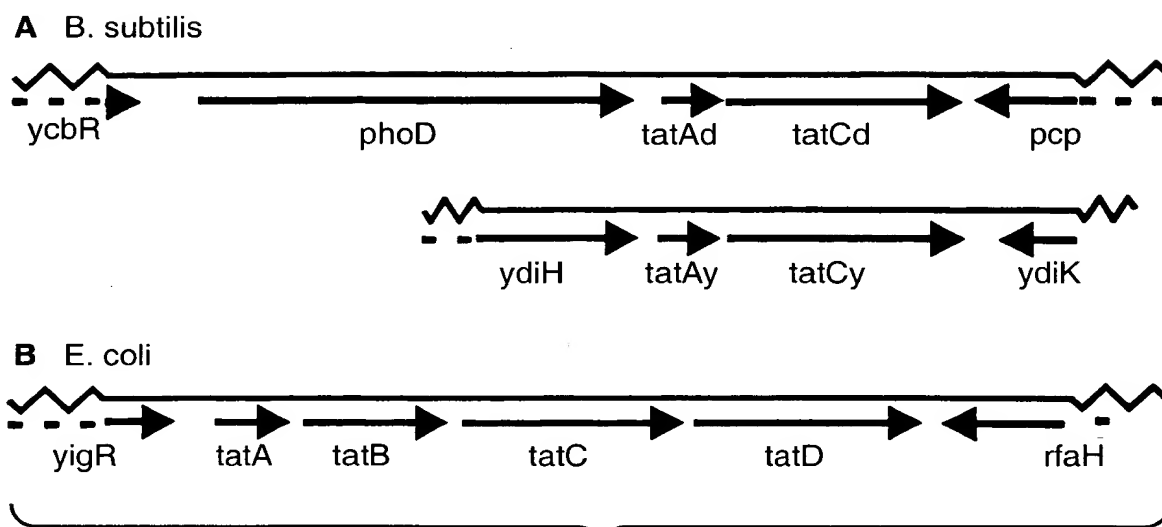
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TatA (Eco)	M- <u>GGISIWQLLII</u> AVIVVLLFGTKKLG-----	26
TatE (Eco)	M- <u>GEISITKLLV</u> VAAALVLLFGTKKLRLR-----	26
TatAy (Bsu)	<u>M</u> --- <u>PIGPGSLAVIA</u> IIVALIIIFGPKKLP-----	25
TatAd (Bsu)	MFSNIGIPGLILIFVIAIIIFGPKSLP-----	27
TatAc (Bsu)	<u>M</u> --- <u>ELSFTKILVIL</u> FVGFVFGPDKLP-----	25
TatB (Eco)	<u>MF</u> - <u>DIGFSELLLVFI</u> IGLVVLGPQRLPVAVKTVAGWIRALRSLATTVQNELTQELKLO	49
	* . . . . . *	
TatA (Eco)	-----SIGSDLGASIKGFKKAMSDDE----PKQDKTSQDADFTAKTI	64
TatE (Eco)	-----TLGGDLGAAIKGFKKAMNDDD----A-AAKKGADVDLQAEKL	63
TatAy (Bsu)	-----ELGKAAGDTLREFKNATKGLT----SDEEEKKKEDQ-----	57
TatAd (Bsu)	-----EIGRAAKRTLLEFKSATKSLV----SGDEKEEKSAELTAVK-	64
TatAc (Bsu)	-----ALGRAAGKALSEFKQATSGLT----QDIRKNDSN-----K-	57
TatB (Eco)	EFQDSLKKVEKASLTNLTPELKASMDLRQAESMKRSYVANDPEKASDEAHTIHP	114
	. . . . . *	
TatA (Eco)	ADKQADTNQE-----QAKTEDAKRHDKEQV	89
TatE (Eco)	SHKE-----	67
TatAy (Bsu)	-----	57
TatAd (Bsu)	-----QDKNAG	70
TatAc (Bsu)	-----EDKQM-	62
TatB (Eco)	VVKDNEAAHEGVTPAAAQTQASSPEQKPETTPEPVVKPAADAEPKTAAPSPSSSDKP	171

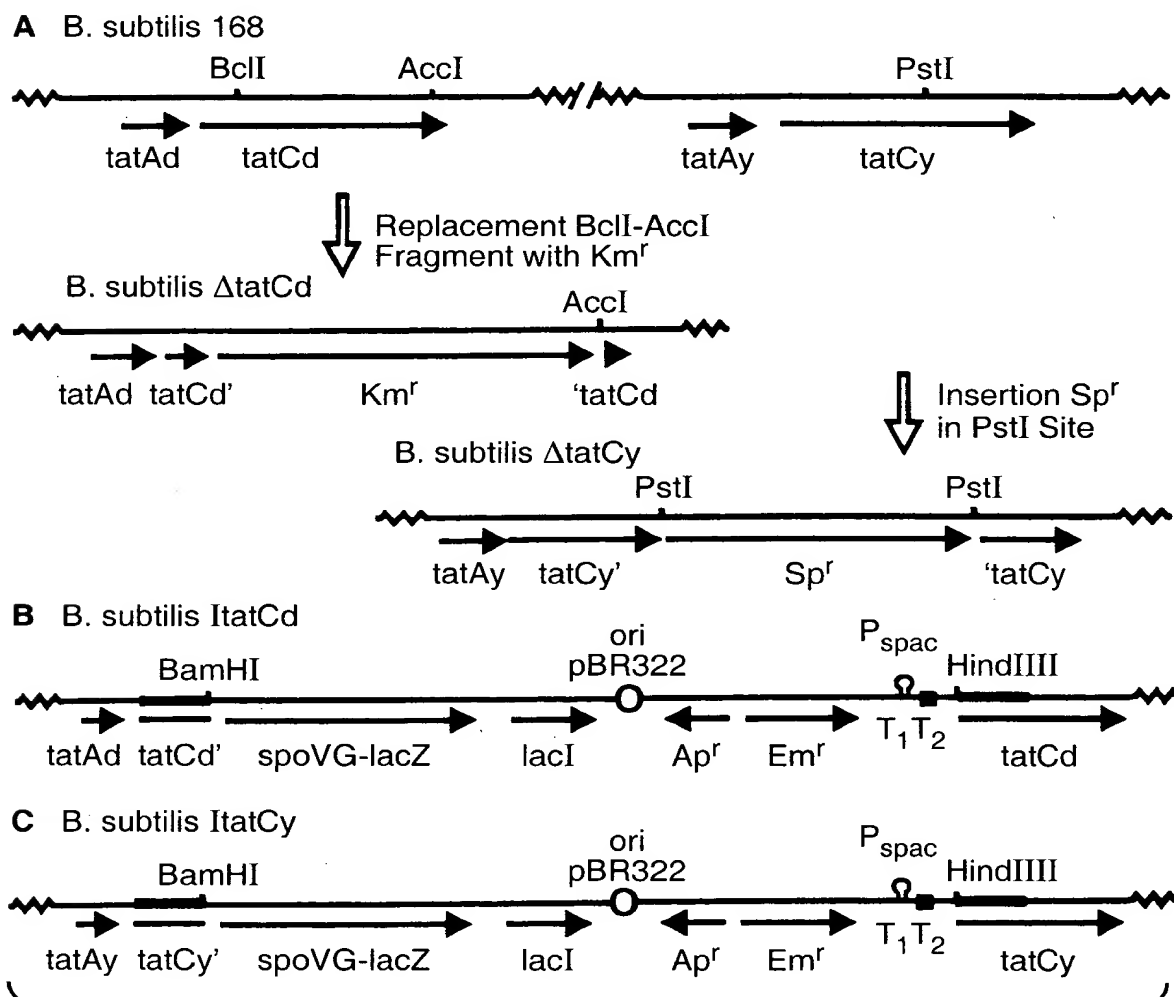
## FIG. 1A

TatC (Eco)	MSVEDTQ--PLITHLIELRK <u>RL</u> LNCCI <del>IAVIVIFLCLVYFANDIYH</del> -LVSAPLIK	51
TatCy (Bsu)	MTRMKVNQMSLLEHIAELRK <u>LL</u> IVALAFVVF <del>FIAGFFLAKPIIVYLQETDEAK</del>	50
TatCd (Bsu)	MDKKETH---LIGHLEELRR <u>RI</u> IVTLAAFFLFLITAF <del>LFVQDIYDWLIRDLDGK</del>	51
	*. . . . * . . . . * . . . . *	
TatC (Eco)	QLPQGSTMATDVASPF <del>FT</del> <u>IKLTFMVSLILSAPVILYQVWAFIAPALYKHERR</u>	105
TatCy (Bsu)	QL----TLNAFNLTDPLYVFMQFAFIIGIVLTSPVILYQ <del>LWAFVSPGLYEKERK</del>	104
TatCd (Bsu)	-----LAVLGPSEILWVY <u>MLSGICAIAASIPVAAYQLWRFVAPALTKTERK</u>	98
	. . . . . * . . . . *	
TatC (Eco)	LVVPLL <del>V</del> ---SSSL <u>LFYIGMAFAYFVVFPLAFGFLANTAPE</u> -GVQVSTD <u>IASYL</u>	155
TatCy (Bsu)	VTLSYI---PVSILLFLAGLSFSYI <del>LPFVVD</del> FMKRISQDLNVNQVIGINEYF	155
TatCd (Bsu)	VTIMYIMYIPGLFALFLAGISFGYFV <del>LPFIVLSFLTHLSSG</del> -HFETMFTADRYF	151
	. . . . . * . . . . *	
TatC (Eco)	<u>SFVMALFMAFGVSFEV</u> PVAIVLLCWMGITSPEDLRKKR <u>PYVLVGAFVVGMLLTP</u>	209
TatCy (Bsu)	HFLQLTIPFGLLFQMPVILMFLTRLGIVTPMFLAKIRK <u>YAYFTLLVIAALITP</u>	209
TatCd (Bsu)	RFMVNL <del>SLPFGFLFEM</del> PLVVMFLTRLGILNPYRLAKA <u>RKLSYFLLIVVSILITP</u>	205
	* . . . . * . . . . * . . . . *	
TatC (Eco)	PDVFSQTL <del>LAIPMYCLFEIGVFFSRE</del> -YVGKGRNREEENDAEAESEKTEE	258
TatCy (Bsu)	PELLSHMMVTVP <del>LLILYEISILISKAAYRKAQKSSAADRDVSSG</del> -----Q	254
TatCd (Bsu)	PDFISDFLVMIP <del>LLVLFEVSVTL</del> SAFVYKKRMRE-----ETAAA-----A	245
	* . . . . * . . . . *	

## FIG. 1B



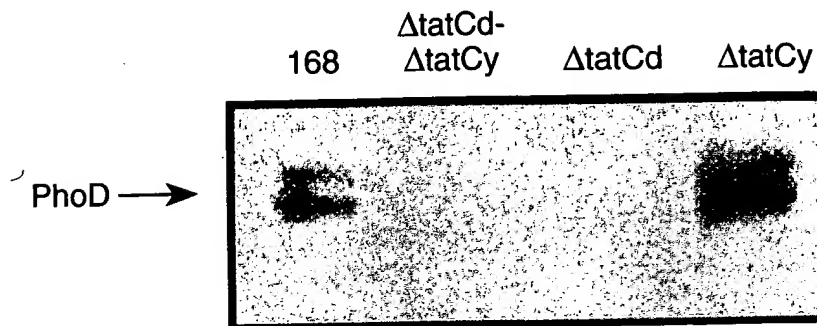
**FIG. 2**



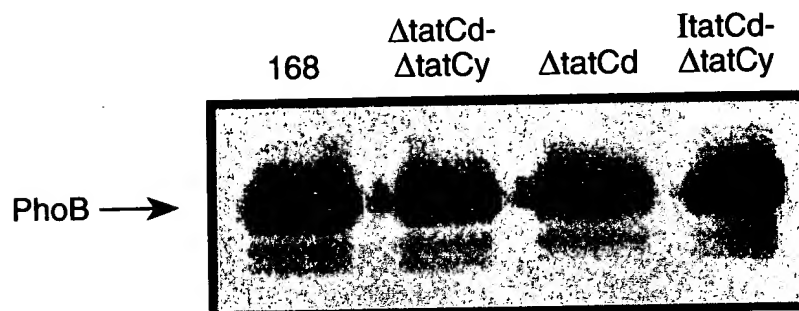
**FIG. 3**



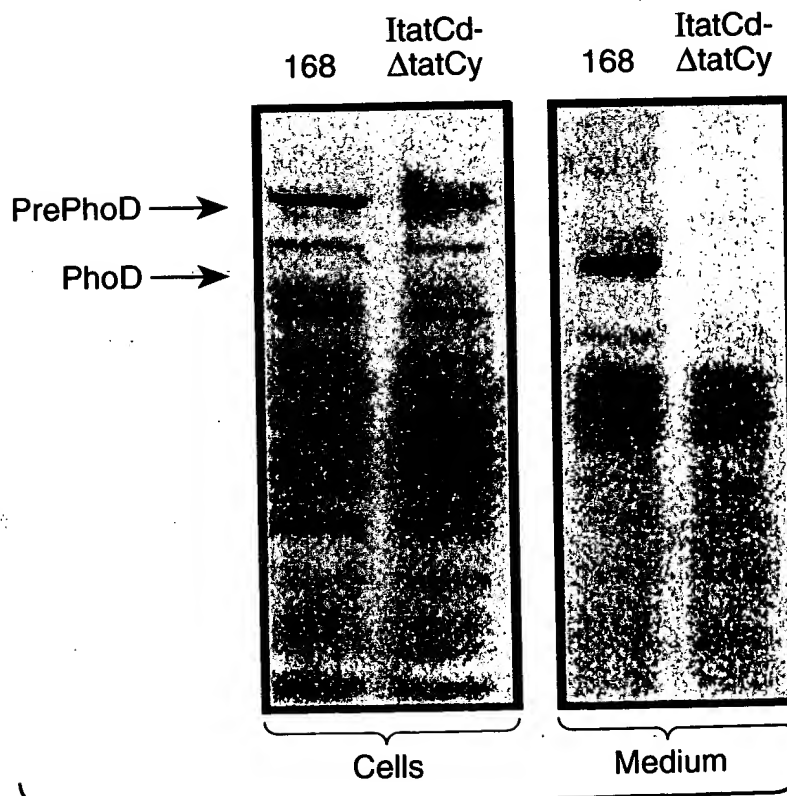
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**FIG. 4A**



**FIG. 4B**



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$\Delta$ tatCd- $\Delta$ tatCy

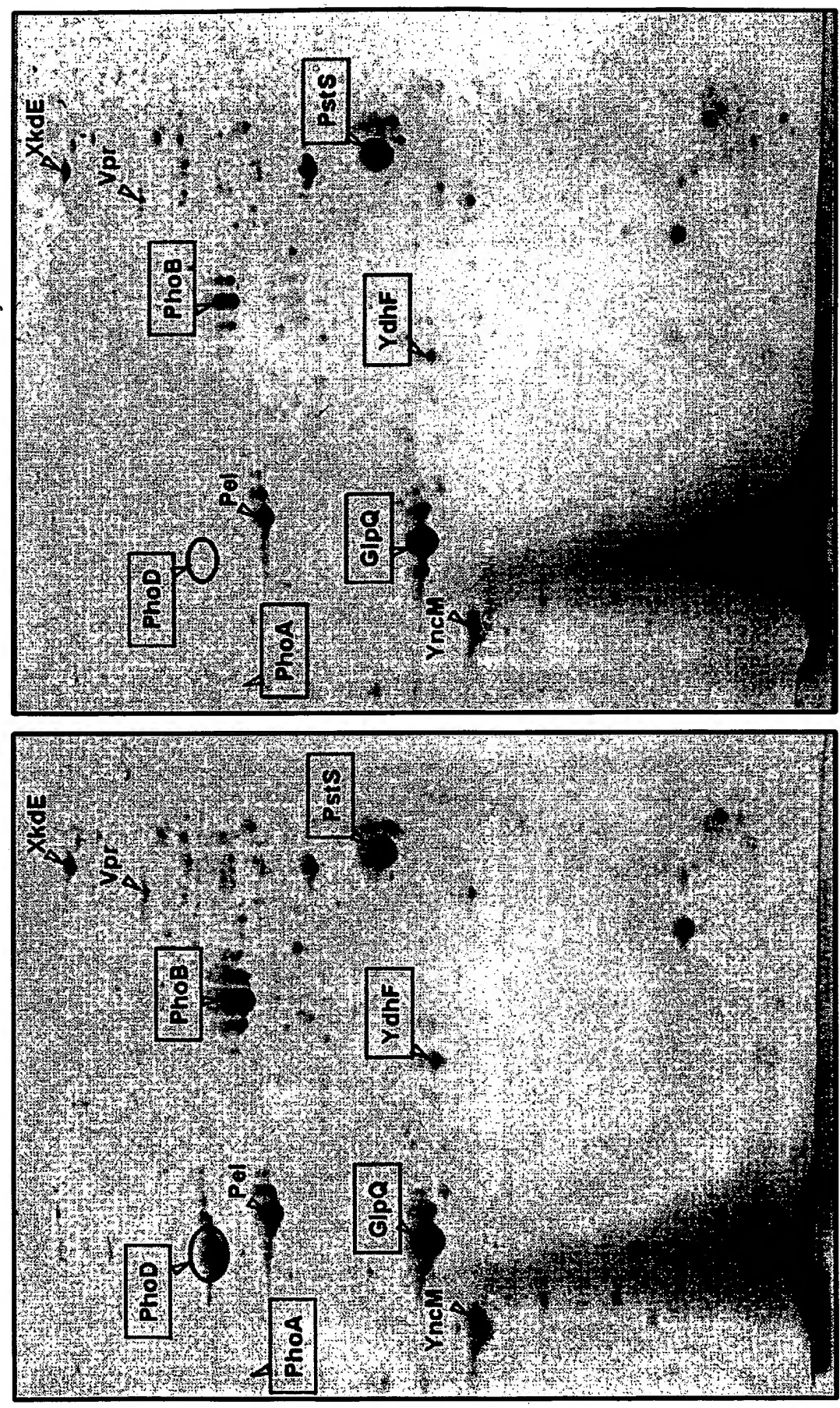
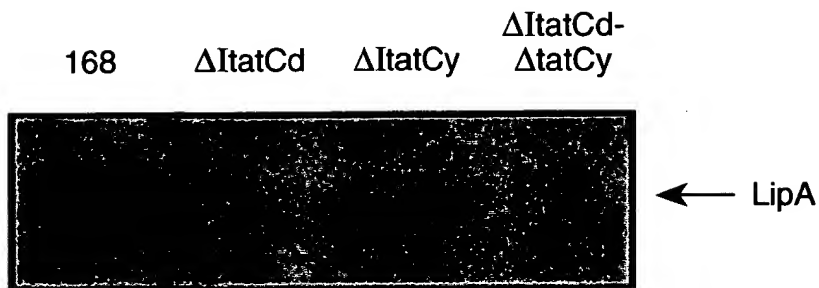


FIG.\_5



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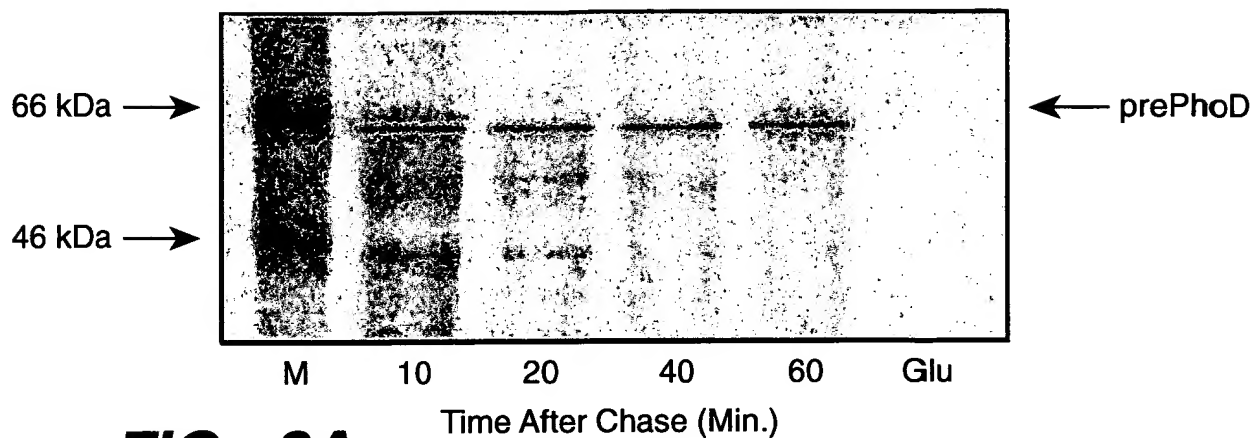
**FIG.\_6**

Protein	N	h	RR-Motif	H	h	C
AlbB	1	0.1	RRILL	27	2.0	AIA
AmyX <sup>TM</sup>	9	-0.8	RRSFE	15	1.1	-
AppB <sup>TM</sup>	8	0.5	RRTLm	19	2.3	-
LipA	7	-1.1	RRIIA	19	1.2	AKA
OppB <sup>TM</sup>	8	-0.6	RRLVY	24	2.0	-
PbpX	2	-2.2	RRRKL	14	2.9	WNA
PhoD	3	-1.3	RRKFI	17	0.9	VGA
QcrA <sup>TM</sup>	1	-1.1	RRQFL	19	1.3	-
TlpA <sup>TM</sup>	1	-0.8	RRLII	21	2.4	-
WapA <sup>W</sup>	1	-3.0	RRNFK	18	2.3	VLA
WprA	8	-1.7	RRKFS	20	1.9	AAA
YceA <sup>TM</sup>	1	-0.4	RRAFI	21	2.2	-
YesM <sup>TM</sup>	1	-1.5	RRMKI	20	2.4	QYA
YesW	1	-1.3	RRSCL	19	2.0	VKA
YfkN <sup>TM</sup>	1	-1.2	RRTHV	17	1.7	IHA
YkpC	8	-1.0	RRVAI	17	2.3	SLA
YkuE	1	-1.3	RRQFL	17	1.0	GYA
YmaC	7	0.0	RRFLL	15	2.4	YSL
YubF <sup>TM</sup>	9	-2.7	RRNTV	23	2.0	-
YuiC	8	0.2	RRLLM	20	1.9	IEA
YvhJ <sup>TM</sup>	2	-1.7	RRKIL	18	2.5	-
YwbN	1	-1.8	RRDIL	23	1.4	QTA

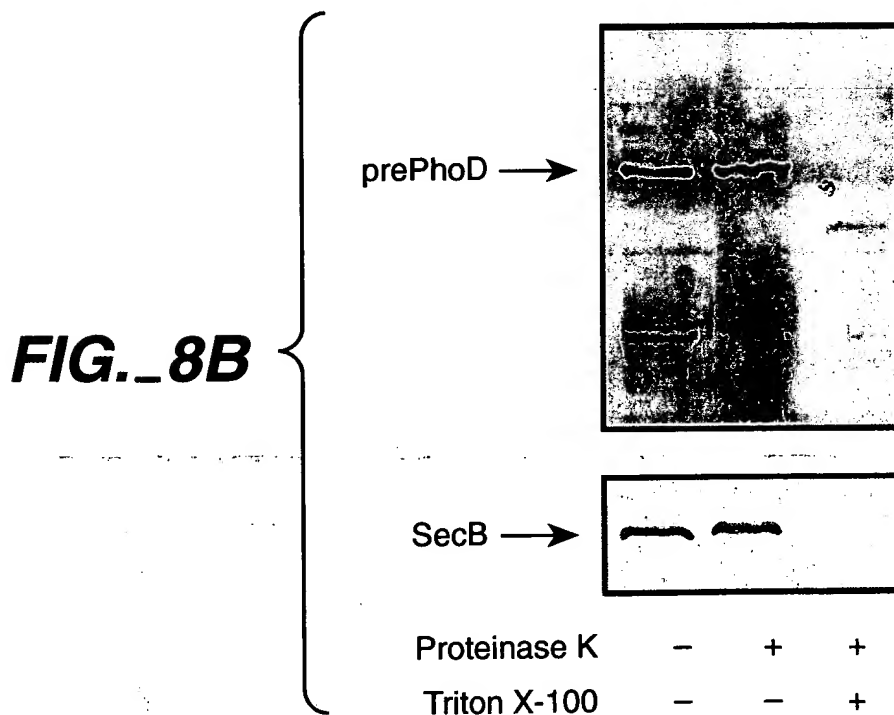
**FIG.\_7**



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**FIG.\_8A**



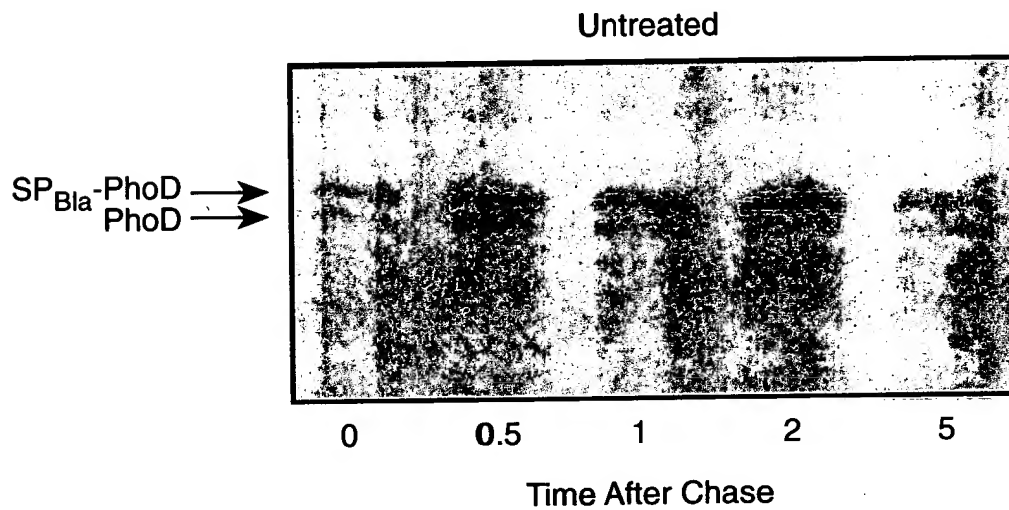
**FIG.\_8B**



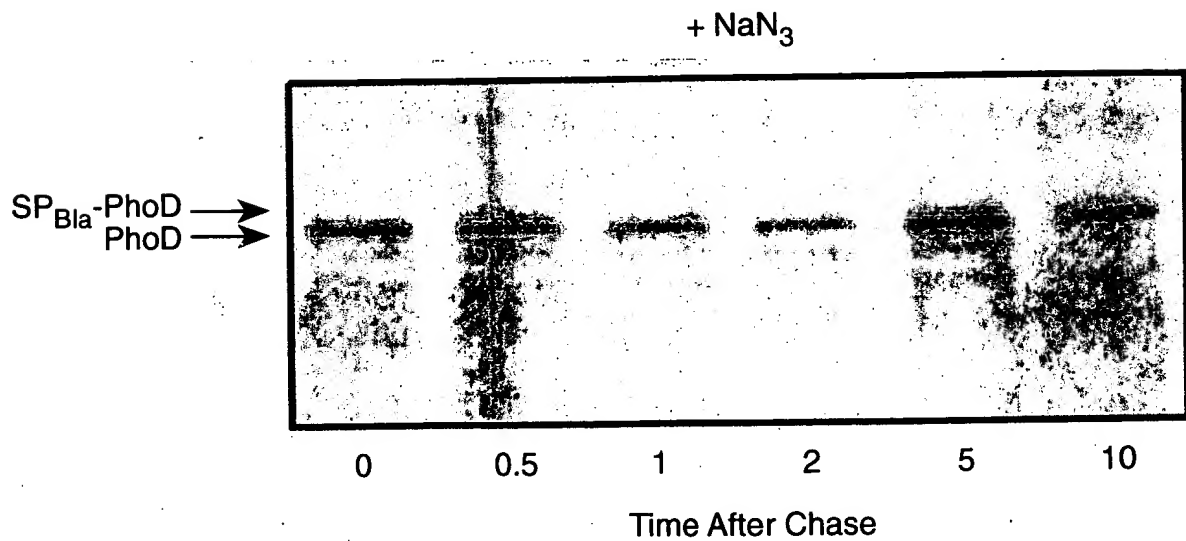
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**FIG.\_9A**

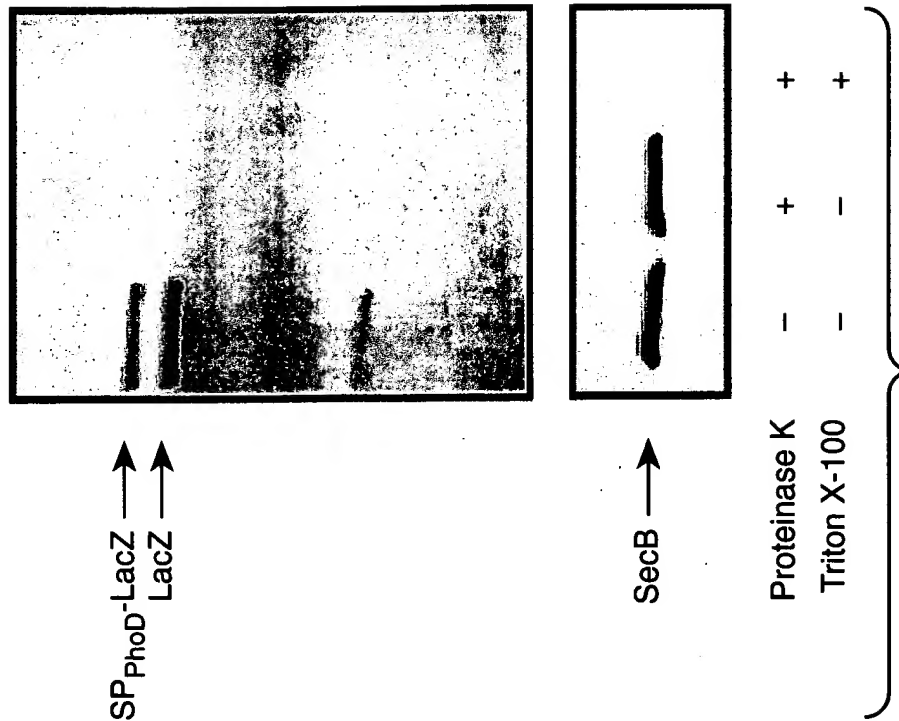


**FIG.\_9B**

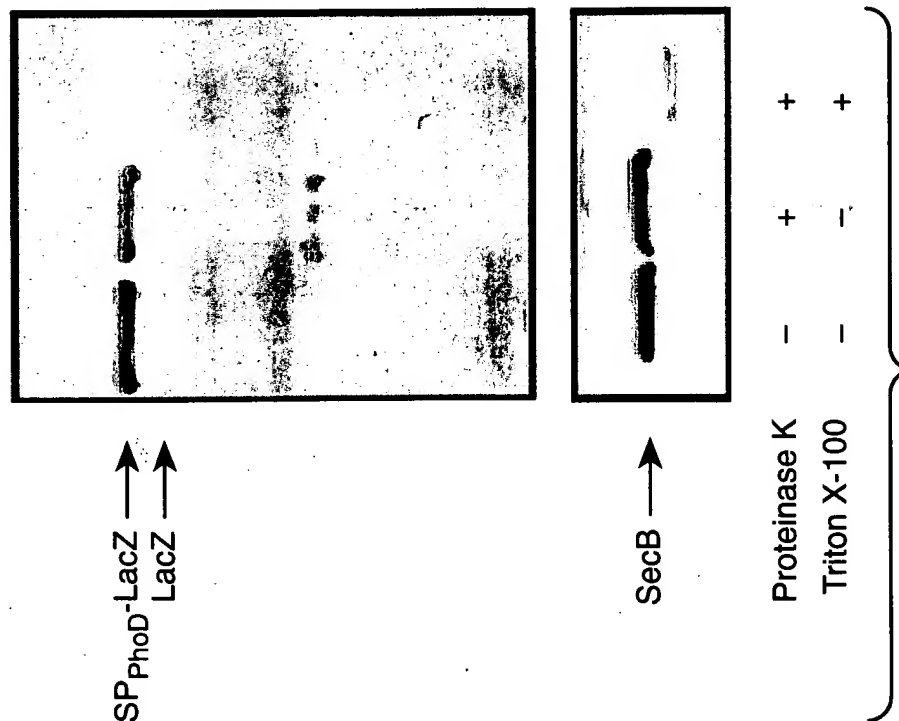


**FIG.\_9C**





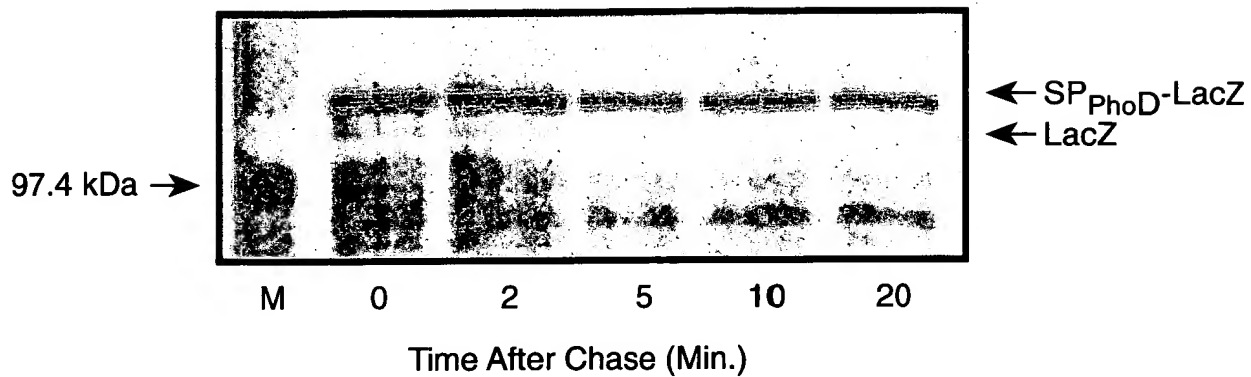
**FIG.\_10B**



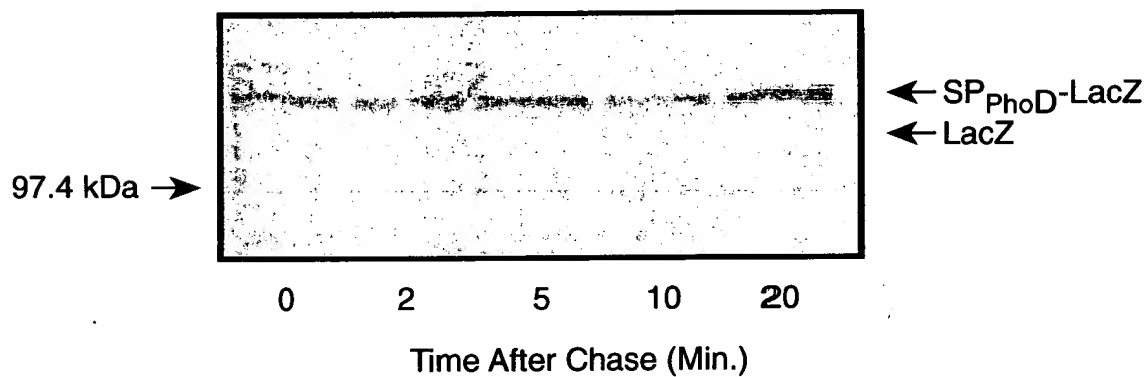
**FIG.\_10A**



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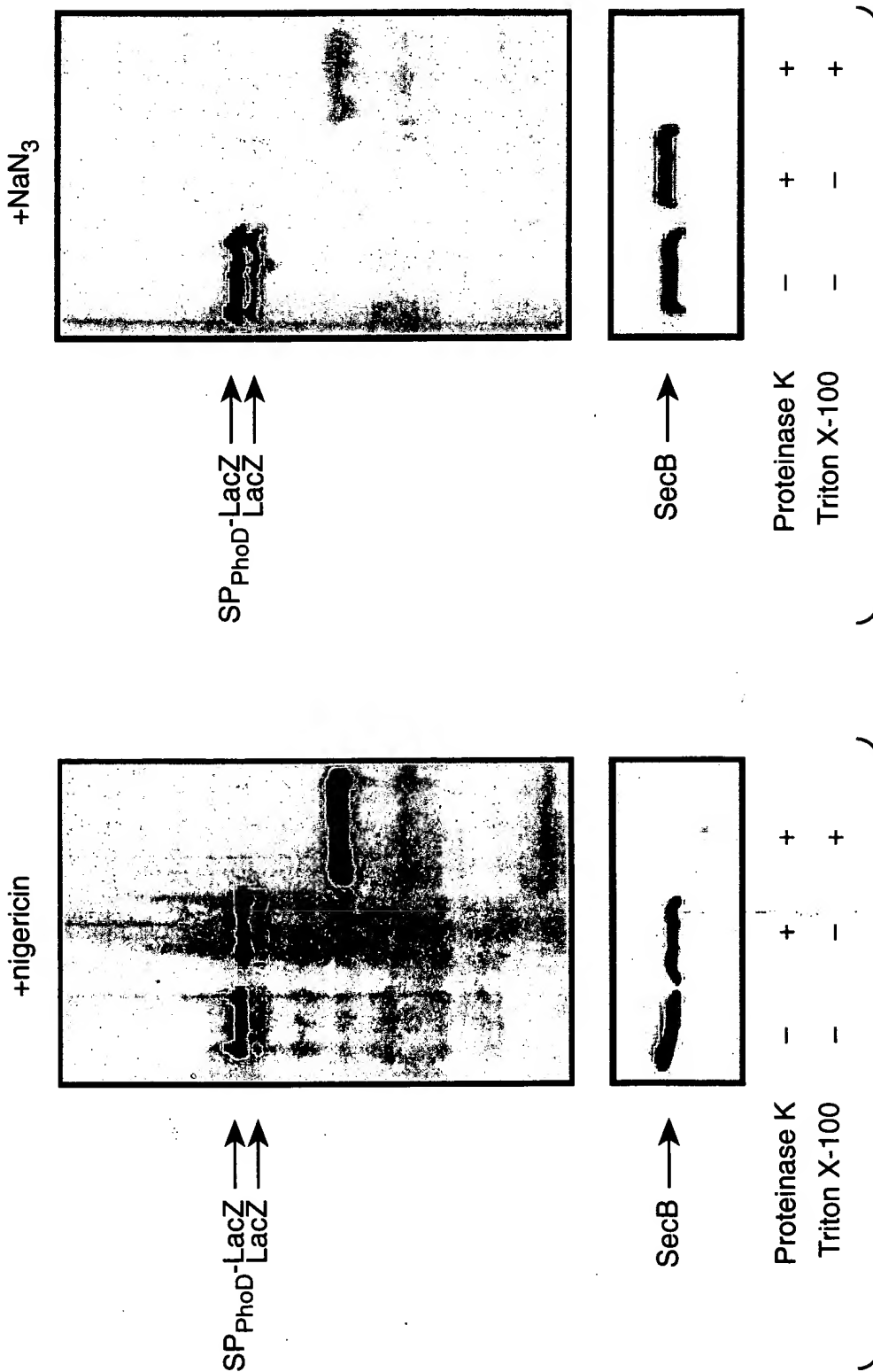
**FIG.\_11A**



**FIG.\_11B**

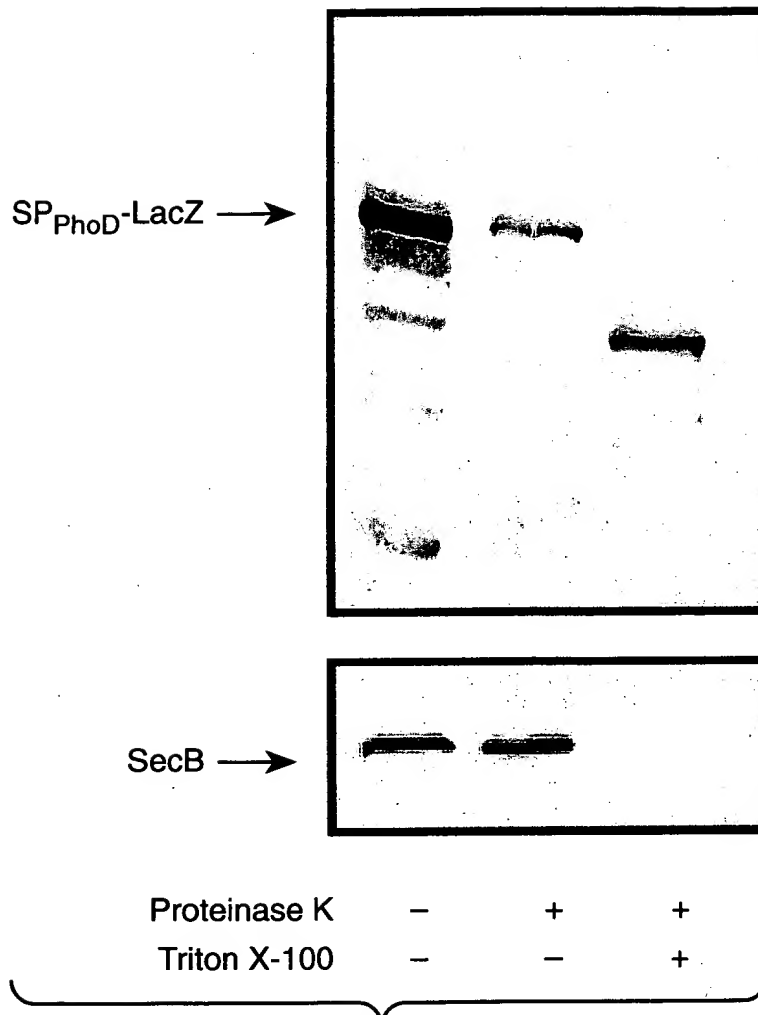


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**FIG. 13**

Homologs in *B. alcalophilus*

TatA

MGGLSVGSVVLIALVALLIFGPKKLPELGKAAGSTLREFKNATK  
GLADDDDDTKSTNVQKEKA

TatC

MTMMTPNQQTSKKKKRKGRKGRVPMQDMSIMDHAEELRRRIF  
VVLAFVIALIGGFFLAVPVITFLQNSPQAADMPFNAFRLTDPLRV  
YMNFAVITALVLIIPVILYQLWAFVSPGLKENEQKATLAYIPIAFL  
LFLAGIAFSYFILLPFVISFMGQMADRLEINEMYGINEYFSFLFQL  
TIPFGLLFQLPVVVMFLTRLGVVTPFTFLRKIRKYAYFALLVIAGII  
TPPELTSHLFVTVPMLILYEISITISAITYRKYHGTTHNGQESAK